

<110> Conklin, Darrell C.
Blumberg, Hal

<130> 97-63C1

<151> 1997-10-06

<151> 1998-10-06

<160> 28

<170> FastSEQ for Windows Version 3.0

 $\langle 210 \rangle$ 1

<211> 876

<212> DNA

<213> Homo sapiens

 $\langle 220 \rangle$

<221> CDS

<222> (119)...(823)

 $\langle 400 \rangle$ 1

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ctggctgagg tcgctggga gctgccgcca gggccaggag gggagcgga cctggaag      118
atg cgc cca ttg gct ggt ggc ctg ctc aag gtg gtg ttc gtg gtc ttc      166
Met Arg Pro Leu Ala Gly Gly Leu Leu Lys Val Val Phe Val Val Phe
      1             5             10             15
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gcc tcc ttg tgt gcc tgg tat tcg ggg tac ctg ctc gca gag ctc att 214
Ala Ser Leu Cys Ala Trp Tyr Ser Gly Tyr Leu Leu Ala Glu Leu Ile
20 25 30

cca gat gca ccc ctg tcc agt gct gcc tat agc atc cgc agc atc ggg	262
Pro Asp Ala Pro Leu Ser Ser Ala Ala Tyr Ser Ile Arg Ser Ile Gly	
35 40 45	
gag agg cct gtc ctc aaa gct cca gtc ccc aaa agg caa aaa tgt gac	310
Glu Arg Pro Val Leu Lys Ala Pro Val Pro Lys Arg Gln Lys Cys Asp	
50 55 60	
cac tgg act ccc tgc cca tct gac acc tat gcc tac agg tta ctc agc	358
His Trp Thr Pro Cys Pro Ser Asp Thr Tyr Ala Tyr Arg Leu Leu Ser	
65 70 75 80	
gga ggt ggc aga agc aag tac gcc aaa atc tgc ttt gag gat aac cta	406
Gly Gly Gly Arg Ser Lys Tyr Ala Lys Ile Cys Phe Glu Asp Asn Leu	
85 90 95	
ctt atg gga gaa cag ctg gga aat gtt gcc aga gga ata aac att gcc	454
Leu Met Gly Glu Gln Leu Gly Asn Val Ala Arg Gly Ile Asn Ile Ala	
100 105 110	
att gtc aac tat gta act ggg aat gtg aca gca aca cga tgt ttt gat	502
Ile Val Asn Tyr Val Thr Gly Asn Val Thr Ala Thr Arg Cys Phe Asp	
115 120 125	
atg tat gaa ggc gat aac tct gga ccg atg aca aag ttt att cag agt	550
Met Tyr Glu Gly Asp Asn Ser Gly Pro Met Thr Lys Phe Ile Gln Ser	
130 135 140	
gct gct cca aaa tcc ctg ctc ttc atg gtg acc tat gac gac gga agc	598
Ala Ala Pro Lys Ser Leu Leu Phe Met Val Thr Tyr Asp Asp Gly Ser	
145 150 155 160	
aca aga ctg aat aac gat gcc aag aat gcc ata gaa gca ctt gga agt	646
Thr Arg Leu Asn Asn Asp Ala Lys Asn Ala Ile Glu Ala Leu Gly Ser	
165 170 175	
aaa gaa atc agg aac atg aaa ttc agg tct agc tgg gta ttt att gca	694
Lys Glu Ile Arg Asn Met Lys Phe Arg Ser Ser Trp Val Phe Ile Ala	
180 185 190	
gca aaa ggc ttg gaa ctc cct tcc gaa att cag aga gaa aag atc aac	742
Ala Lys Gly Leu Glu Leu Pro Ser Glu Ile Gln Arg Glu Lys Ile Asn	
195 200 205	

10039376.103664

cag ata gaa ggc tgc ata ccc aaa gaa cga agc tgacactgca gggctctgag 843
Gln Ile Glu Gly Cys Ile Pro Lys Glu Arg Ser
225 230 235

taaatgtgtt ctgtataaac aaatgcagct gga 876

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<210> 2
<211> 235
<212> PRT
<213> Homo sapiens
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Pro	Asp	Ala	Pro	Leu	Ser	Ser	Ala	Ala	Tyr	Ser	Ile	Arg	Ser	Ile	Gly	
		35					40					45				
Glu	Arg	Pro	Val	Leu	Lys	Ala	Pro	Val	Pro	Lys	Arg	Gln	Lys	Cys	Asp	
	50					55					60					
His	Trp	Thr	Pro	Cys	Pro	Ser	Asp	Thr	Tyr	Ala	Tyr	Arg	Leu	Leu	Ser	
65					70					75					80	
Gly	Gly	Gly	Arg	Ser	Lys	Tyr	Ala	Lys	Ile	Cys	Phe	Glu	Asp	Asn	Leu	
				85					90						95	
Leu	Met	Gly	Glu	Gln	Leu	Gly	Asn	Val	Ala	Arg	Gly	Ile	Asn	Ile	Ala	
			100					105						110		
Ile	Val	Asn	Tyr	Val	Thr	Gly	Asn	Val	Thr	Ala	Thr	Arg	Cys	Phe	Asp	
		115					120					125				
Met	Tyr	Glu	Gly	Asp	Asn	Ser	Gly	Pro	Met	Thr	Lys	Phe	Ile	Gln	Ser	
	130					135					140					
Ala	Ala	Pro	Lys	Ser	Leu	Leu	Phe	Met	Val	Thr	Tyr	Asp	Asp	Gly	Ser	
145					150					155					160	
Thr	Arg	Leu	Asn	Asn	Asp	Ala	Lys	Asn	Ala	Ile	Glu	Ala	Leu	Gly	Ser	
				165					170						175	
Lys	Glu	Ile	Arg	Asn	Met	Lys	Phe	Arg	Ser	Ser	Trp	Val	Phe	Ile	Ala	
			180					185					190			
Ala	Lys	Gly	Leu	Glu	Leu	Pro	Ser	Glu	Ile	Gln	Arg	Glu	Lys	Ile	Asn	
	195					200						205				

His Ser Asp Ala Lys Asn Asn Arg Tyr Ser Gly Trp Pro Ala Glu Ile
 210 215 220
 Gln Ile Glu Gly Cys Ile Pro Lys Glu Arg Ser
 225 230 235

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 <223> Motif 1. corresponding to residues 127 to 129 of
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<210> 4
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<400> 4
 Tyr Asp Asp
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<210> 5
 <211> 3
 <212> PRT
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<220>
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 SEQ ID NO:2

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 Leu Gly Ser
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10039876-103601

<210> 6
 <211> 3
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<220>
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<400> 6
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<210> 7
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 <212> PRT
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<210> 8
 <211> 705
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<221> variation
 <222> (1)...(705)
 <223> N is any nucleotide

<221> misc_feature
 <222> (1)...(705)
 <223> n = A,T,C or G

10039276 "102601
 222601

<400> 8

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gcntggtayw snggntayyt nytnngcgar ytnathccng aygcncnyt nwsnwsngcn	120
gcntaywsna thmgwnsnat hggngarmgn ccngtnytna argcncngt nccnaarmgn	180
caraartgyg aycaytgac nccntgyccn wsngayacnt aygcntaymg nytnytnwsn	240
ggngngngnm gnwsnaarta ygcnaarath tgyttygarg ayaayytnyt natggngar	300
carytnggna aygtngcnmg nggnathaay athgcnathg tnaaytaygt nacnggnaay	360
gtnacngcna cnmgntgytt ygayatgtay gargngaya aywsnggncc natgacnaar	420
ttyathcarw sngcngcncc naarwsnytn ytnttyatgg tnacntayga ygaygnwsn	480
acnmgnytna ayaaygaygc naaraaygcn athgargcny tnggnwsnaa rgarathmgn	540
aayatgaart tymgnwsnws ntgggtntty athgngcna arggnytna rytncnwsn	600
garathcarm gngaraarat haaycaywsn gaygcnaara ayaaymgnta ywsnggntgg	660
ccngngara thcarathga rggntgyath ccnaargarm gnwsn	705

<210> 9

<211> 19

<212> DNA

<213> Artificial Sequence

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<223> Oligonucleotide primer ZC695

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<210> 10

<211> 26

<212> DNA

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<223> Oligonucleotide primer ZC7231

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tttttttttt tttttttttt tttttv	26
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<210> 11

<211> 20

<212> DNA

<213> Artificial Sequence

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<223> Oligonucleotide primer ZC13695

1003929.103604

<400> 11
cccttccgaa attcagagag 20

<210> 12
<211> 20
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<223> Oligonucleotide primer ZC13789

<400> 12
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<400> 13
ccagctgttc tcccataagt 20

<210> 14
<211> 21
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<220>
<223> Oligonucleotide primer ZC14069

<400> 14
cttgcatcg ttattcagtc t 21

<210> 15
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<223> Oligonucleotide primer ZC13072

<400> 15

aggtcctggg caagtgcctgc

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<210> 16

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Oligonucleotide primer ZC13956

<400> 16

gtggtgttcg tggctctc

18

<210> 17

<211> 18

<212> DNA

<213> Artificial Sequence

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<223> Oligonucleotide primer ZC13957

<400> 17

cgatgctgcg gatgctat

18

<210> 18

<211> 243

<212> PRT

<213> Artificial Sequence

<220>

<223> z219aCEE polypeptide with C-terminal GluGlu tag

<400> 18

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			20					25					30		
Pro	Asp	Ala	Pro	Leu	Ser	Ser	Ala	Ala	Tyr	Ser	Ile	Arg	Ser	Ile	Gly
			35				40						45		

10039376.10039376.1

Glu Arg Pro Val Leu Lys Ala Pro Val Pro Lys Arg Gln Lys Cys Asp
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 His Trp Thr Pro Cys Pro Ser Asp Thr Tyr Ala Tyr Arg Leu Leu Ser
 65 70 75 80
 Gly Gly Gly Arg Ser Lys Tyr Ala Lys Ile Cys Phe Glu Asp Asn Leu
 85 90 95
 Leu Met Gly Glu Gln Leu Gly Asn Val Ala Arg Gly Ile Asn Ile Ala
 100 105 110
 Ile Val Asn Tyr Val Thr Gly Asn Val Thr Ala Thr Arg Cys Phe Asp
 115 120 125
 Met Tyr Glu Gly Asp Asn Ser Gly Pro Met Thr Lys Phe Ile Gln Ser
 130 135 140
 Ala Ala Pro Lys Ser Leu Leu Phe Met Val Thr Tyr Asp Asp Gly Ser
 145 150 155 160
 Thr Arg Leu Asn Asn Asp Ala Lys Asn Ala Ile Glu Ala Leu Gly Ser
 165 170 175
 Lys Glu Ile Arg Asn Met Lys Phe Arg Ser Ser Trp Val Phe Ile Ala
 180 185 190
 Ala Lys Gly Leu Glu Leu Pro Ser Glu Ile Gln Arg Glu Lys Ile Asn
 195 200 205
 His Ser Asp Ala Lys Asn Asn Arg Tyr Ser Gly Trp Pro Ala Glu Ile
 210 215 220
 Gln Ile Glu Gly Cys Ile Pro Lys Glu Arg Ser Gly Ser Glu Tyr Met
 225 230 235 240
 Pro Met Glu

<210> 19

<211> 25

<212> DNA

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<223> Oligonucleotide primer ZC14870

<400> 19

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25

<210> 20

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<212> DNA

<213> Artificial Sequence

1003936.10001

<220>

<223> Oligonucleotide primer ZC15101

<400> 20

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24

<210> 21

<211> 8

<212> PRT

<213> Artificial Sequence

<220>

<223> Glu-Glu tag sequence with linker

<400> 21

Gly Ser Glu Tyr Met Pro Met Glu

1

5

<210> 22

<211> 20

<212> DNA

<213> Artificial Sequence

<220>

<223> Oligonucleotide primer ZC13006

<400> 22

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<210> 23

<211> 19

<212> DNA

<213> Artificial Sequence

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<223> Oligonucleotide primer ZC13007

<400> 23

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19

<210> 24

<211> 6

<212> PRT

<223> EE peptide sequence

Glu Tyr Met Pro Val Asp
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<213> Artificial Sequence

<223> Oligonucleotide primer ZC12700

21

<213> Artificial Sequence

<223> Oligonucleotide primer ZC12742

20

<213> Artificial Sequence

<223> Oligonucleotide primer ZC17184

32

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32